

**In the Claims**

1. (Original) A pharmaceutical composition comprising:

an aqueous carrier;

from 0.1 mg/ml to 20 mg/ml of the composition of a pharmaceutically acceptable salt of

a) a peptide comprising at least 12 and at most 30 consecutive amino acids having a sequence corresponding to

(i) a sequence of amino acids found within a complementarity-determining region (CDR) of a heavy or a light chain of a human monoclonal anti-DNA 16/6 Id antibody, or

(ii) a sequence of amino acids found within a complementarity-determining region (CDR) of a heavy or a light chain of a pathogenic anti-DNA monoclonal antibody that induces a systemic lupus erythematosus (SLE)-like disease response in mice, or

b) a peptide comprising consecutive amino acids having the sequence

(i) TGYX<sub>1</sub>X<sub>2</sub>X<sub>3</sub>X<sub>4</sub>X<sub>5</sub>QSPEKSLEWIG (SEQ ID NO:11)

wherein X<sub>1</sub> is Met, Ala or Val; X<sub>2</sub> is Gln, Asp, Glu or Arg; X<sub>3</sub> is Trp or Ala; X<sub>4</sub> is Val or Ser; and X<sub>5</sub> is Lys, Glu or Ala;

(ii) EINPSTGGX<sub>6</sub>X<sub>7</sub>X<sub>8</sub>X<sub>9</sub>X<sub>10</sub>X<sub>11</sub>X<sub>12</sub>KAKAT (SEQ ID NO:12)

wherein X<sub>6</sub> and X<sub>7</sub> are each Thr, Val or Ala; X<sub>8</sub> is Tyr or Phe; X<sub>9</sub> is Asn or Asp; X<sub>10</sub> is Gln or Glu; X<sub>11</sub> is Lys or Glu, and X<sub>12</sub> is Phe or Tyr;

(iii) YYCARX<sub>13</sub>X<sub>14</sub>X<sub>15</sub>X<sub>16</sub>PYAX<sub>17</sub>X<sub>18</sub>YWGQGS (SEQ ID NO:13)

wherein X<sub>13</sub> is Phe, Thr or Gly; X<sub>14</sub> is Leu, Ala or

Ser; X<sub>15</sub> is Trp or Ala; X<sub>16</sub> is Glu or Lys; X<sub>17</sub> is Met or Ala, and X<sub>18</sub> is Asp, Lys or Ser;

(iv) GYNX<sub>19</sub>X<sub>20</sub>X<sub>21</sub>X<sub>22</sub>X<sub>23</sub>X<sub>24</sub>SHGX<sub>25</sub>X<sub>26</sub>LEWIG (SEQ ID NO:14)  
wherein X<sub>19</sub> is Met or Ala; X<sub>20</sub> is Asn, Asp or Arg; X<sub>21</sub> is Trp or Ala; X<sub>22</sub> is Val or Ser; X<sub>23</sub> is Lys or Glu; X<sub>24</sub> is Gln or Ala; X<sub>25</sub> is Lys or Glu, and X<sub>26</sub> is Ser or Ala;

(v) YYCARX<sub>27</sub>X<sub>28</sub>X<sub>29</sub>YGX<sub>30</sub>X<sub>31</sub>X<sub>32</sub>GQTL (SEQ ID NO:15)  
wherein X<sub>27</sub> is Ser or Phe; X<sub>28</sub> is Gly or Ala; X<sub>29</sub> is Arg, Ala or Glu; X<sub>30</sub> is Asn or Asp; X<sub>31</sub> is Tyr or Phe, and X<sub>32</sub> is Trp, His or Ala;

(vi) X<sub>33</sub>YYWSWIX<sub>34</sub>QX<sub>35</sub>PX<sub>36</sub>X<sub>37</sub>GX<sub>38</sub>EWIG (SEQ ID NO:16)  
wherein X<sub>33</sub> is Gly or Thr Gly; X<sub>34</sub> is Arg or Lys; X<sub>35</sub> is Pro or Ser; X<sub>36</sub> is Gly or Glu; X<sub>37</sub> is Lys or Asp; and X<sub>38</sub> is Glu, Leu or Ser;

(vii) YYCARX<sub>39</sub>LLX<sub>40</sub>X<sub>41</sub>X<sub>42</sub>X<sub>43</sub>X<sub>44</sub>DVDYX<sub>45</sub>GX<sub>46</sub>DV (SEQ ID NO:17)  
wherein X<sub>39</sub> is Gly or Phe; X<sub>40</sub> is Arg or Ala; X<sub>41</sub> is Gly or Ala; X<sub>42</sub> is Gly or Ala; X<sub>43</sub> is Trp or Ala; X<sub>44</sub> is Asn or Ala; X<sub>45</sub> is Tyr or Trp; X<sub>46</sub> is Met or Gln;

(viii) FSGYYWS (SEQ ID NO:8);

(ix) EINHSGSTNYKTSLS (SEQ ID NO:9); or

(x) GLLRGGWNDVDYYYGMDV (SEQ ID NO:10), or

- c) a peptide comprising consecutive amino acids having a sequence of any of a) and b), or at least two of the sequences in (a)(i), (a)(ii) and (b)(i) through (b)(x), or
- d) a peptide comprising consecutive amino acids having a sequence comprising at least two identical sequences included in (a)(i), (a)(ii) and (b)(i) through (b)(x); and

a solubility enhancer selected from the group consisting of dimethyl-acetamide, polyethylene glycol, polyoxylated castor

oil, N-methyl-2-pyrrolidinone, 1-ethenyl-2-pyrrolidinone, polyoxyethylene sorbitan esters, and a substituted  $\beta$ -cyclodextrin,

wherein both the peptide and the solubility enhancer are dissolved in the aqueous carrier; and

wherein the composition has a pH between 4 and 9.

2. (Original) The pharmaceutical composition of claim 1, wherein at least 0.5 mg/ml of the composition is the pharmaceutically acceptable salt of the peptide.
3. (Previously Presented) The pharmaceutical composition of claim 1, wherein the peptide has a sequence selected from the group consisting of:

NH<sub>2</sub>- Thr Gly Tyr Tyr Met Gln Trp Val Lys Gln Ser Pro Glu Lys Ser  
Leu Glu-Trp Ile Gly-COOH (SEQ ID NO:1);

NH<sub>2</sub>- Glu Ile Asn Pro Ser Thr Gly Gly Thr Thr Tyr Asn Gln Lys Phe  
Lys Ala Lys Ala Thr-COOH (SEQ ID NO:2);

NH<sub>2</sub>- Tyr Tyr Cys Ala Arg Phe Leu Trp Glu Pro Tyr Ala Met Asp Tyr  
Trp Gly Gln Gly Ser-COOH (SEQ ID NO:3);

NH<sub>2</sub>- Gly Tyr Asn Met Asn Trp Val Lys Gln Ser His Gly Lys Ser Leu  
Glu Trp Ile Gly-COOH (SEQ ID NO:4);

NH<sub>2</sub>- Tyr Tyr Cys Ala Arg Ser Gly Arg Tyr Gly Asn Tyr Trp Gly Gln  
Thr Leu -COOH (SEQ ID NO:5);

NH<sub>2</sub>-Gly Tyr Tyr Trp Ser Trp Ile Arg Gln Pro Pro Gly Lys Gly Glu  
Glu Trp Ile Gly-COOH (SEQ ID NO:6);

NH<sub>2</sub>-Tyr Tyr Cys Ala Arg Gly Leu Leu Arg Gly Gly Trp Asn Asp Val  
Asp Tyr Tyr Gly Met Asp Val-COOH (SEQ ID NO:7);

NH<sub>2</sub>- Phe Ser Gly Tyr Tyr Trp Ser-COOH (SEQ ID NO:8);

NH<sub>2</sub>- Glu Ile Asn His Ser Gly Ser Thr Asn Tyr Lys Thr Ser Leu Lys  
Ser-COOH (SEQ ID NO:9); and

NH<sub>2</sub>- Gly Leu Leu Arg Gly Gly Trp Asn Asp Val Asp Tyr Tyr Tyr Gly

Met Asp Val-COOH (SEQ ID NO:10).

4. (Original) The pharmaceutical composition of claim 1, wherein the peptide comprises consecutive amino acids having the sequence

X<sub>33</sub>YYWSWIX<sub>34</sub>QX<sub>35</sub>PX<sub>36</sub>X<sub>37</sub>GX<sub>38</sub>EWIG (SEQ ID NO:16)

wherein X<sub>33</sub> is Gly or Thr Gly; X<sub>34</sub> is Arg or Lys; X<sub>35</sub> is Pro or Ser; X<sub>36</sub> is Gly or Glu; X<sub>37</sub> is Lys or Asp; and X<sub>38</sub> is Glu, Leu or Ser.

5. (Previously Presented) The pharmaceutical composition of claim 1, wherein the solubility enhancer is a substituted  $\beta$ -cyclodextrin.
6. (Original) The pharmaceutical composition of claim 5, wherein the substituted  $\beta$ -cyclodextrin is a hydroxypropyl, a sulfobutyl ether, or asulfopropyl ether substituted  $\beta$ -cyclodextrin.
7. (Original) The pharmaceutical composition of claim 6, wherein the substituted  $\beta$ -cyclodextrin is a substituted sulfobutyl ether  $\beta$ -cyclodextrin.
8. (Previously Presented) The pharmaceutical composition of claim 1, wherein the concentration of peptide in solution is at least 1 mg/ml.
9. (Previously Presented) The pharmaceutical composition of claim 1, wherein the concentration of peptide in solution is at least 2.5 mg/ml.
10. (Previously Presented) The pharmaceutical composition of claim 1, wherein the composition has a pH between 6.5 and 8.5.

11. (Original) The pharmaceutical composition of claim 10, wherein the composition has a pH between 7.5 and 8.5.
12. (Previously Presented) The pharmaceutical composition of claim 1, wherein the pharmaceutically acceptable salt is an acetate salt.
13. (Original) The pharmaceutical composition of claim 5, wherein the pharmaceutically acceptable salt is an acetate salt, and the substituted  $\beta$ -cyclodextrin is hepta-(sulfobutyl ether)- $\beta$ -cyclodextrin.
14. (Withdrawn) A method of alleviating symptoms of systemic lupus erythematosus (SLE) in a human subject comprising administering to the human subject the pharmaceutical composition of claim 1 in an amount effective to alleviate the symptoms of the SLE in the human subject.
15. (Canceled)
16. (Withdrawn) A process for manufacturing the pharmaceutical composition of claim 1, comprising the steps of:
  - a) preparing a solution of dimethyl-acetamide, polyethylene glycol, polyoxylated castor oil, N-methyl-2-pyrrolidinone, 1-ethenyl-2-pyrrolidinone, polyoxyethylene sorbitan esters, or a substituted  $\beta$ -cyclodextrin in an aqueous carrier at a predetermined concentration;
  - b) adding a predetermined amount of a pharmaceutically acceptable salt of
    - 1) a peptide comprising at least 12 and at most 30 consecutive amino acids having a sequence corresponding

to

- (i) a sequence of amino acids found within a complementarity-determining region (CDR) of a heavy or a light chain of a human monoclonal anti-DNA 16/6 Id antibody, or
  - (ii) a sequence of amino acids found within a complementarity-determining region (CDR) of a heavy or a light chain of a pathogenic anti-DNA monoclonal antibody that induces a systemic lupus erythematosus (SLE)-like disease response in mice,
- 2) a peptide comprising amino acids having the sequence
- (i) TGYX<sub>1</sub>X<sub>2</sub>X<sub>3</sub>X<sub>4</sub>X<sub>5</sub>QSPEKSLEWIG (SEQ ID NO:11)  
wherein X<sub>1</sub> is Met, Ala or Val; X<sub>2</sub> is Gln, Asp, Glu or Arg; X<sub>3</sub> is Trp or Ala; X<sub>4</sub> is Val or Ser; and X<sub>5</sub> is Lys, Glu or Ala;
  - (ii) EINPSTGGX<sub>6</sub>X<sub>7</sub>X<sub>8</sub>X<sub>9</sub>X<sub>10</sub>X<sub>11</sub>X<sub>12</sub>KAKAT (SEQ ID NO:12)  
wherein X<sub>6</sub> and X<sub>7</sub> are each Thr, Val or Ala; X<sub>8</sub> is Tyr or Phe; X<sub>9</sub> is Asn or Asp; X<sub>10</sub> is Gln or Glu; X<sub>11</sub> is Lys or Glu, and X<sub>12</sub> is Phe or Tyr;
  - (iii) YYCARX<sub>13</sub>X<sub>14</sub>X<sub>15</sub>X<sub>16</sub>PYAX<sub>17</sub>X<sub>18</sub>YWGQGS (SEQ ID NO:13)  
wherein X<sub>13</sub> is Phe, Thr or Gly; X<sub>14</sub> is Leu, Ala or Ser; X<sub>15</sub> is Trp or Ala; X<sub>16</sub> is Glu or Lys; X<sub>17</sub> is Met or Ala, and X<sub>18</sub> is Asp, Lys or Ser;
  - (iv) GYNX<sub>19</sub>X<sub>20</sub>X<sub>21</sub>X<sub>22</sub>X<sub>23</sub>X<sub>24</sub>SHGX<sub>25</sub>X<sub>26</sub>LEWIG (SEQ ID NO:14)  
wherein X<sub>19</sub> is Met or Ala; X<sub>20</sub> is Asn, Asp or Arg; X<sub>21</sub> is Trp or Ala; X<sub>22</sub> is Val or Ser; X<sub>23</sub> is Lys or Glu; X<sub>24</sub> is Gln or Ala; X<sub>25</sub> is Lys or Glu, and X<sub>26</sub> is Ser or Ala;
  - (v) YYCARX<sub>27</sub>X<sub>28</sub>X<sub>29</sub>YGX<sub>30</sub>X<sub>31</sub>X<sub>32</sub>GQTL (SEQ ID NO:15)  
wherein X<sub>27</sub> is Ser or Phe; X<sub>28</sub> is Gly or Ala; X<sub>29</sub> is Arg, Ala or Glu; X<sub>30</sub> is Asn or Asp; X<sub>31</sub> is Tyr or Phe, and X<sub>32</sub> is Trp, His or Ala;

(vi) X<sub>33</sub>YYWSWIX<sub>34</sub>QX<sub>35</sub>PX<sub>36</sub>X<sub>37</sub>GX<sub>38</sub>EWIG (SEQ ID NO:16)

wherein X<sub>33</sub> is Gly or Thr Gly; X<sub>34</sub> is Arg or Lys; X<sub>35</sub> is Pro or Ser; X<sub>36</sub> is Gly or Glu; X<sub>37</sub> is Lys or Asp; and X<sub>38</sub> is Glu, Leu or Ser;

(vii)YYCARX<sub>39</sub>LLX<sub>40</sub>X<sub>41</sub>X<sub>42</sub>X<sub>43</sub>X<sub>44</sub>DVDYX<sub>45</sub>GX<sub>46</sub>DV (SEQ ID NO:17)

wherein X<sub>39</sub> is Gly or Phe; X<sub>40</sub> is Arg or Ala; X<sub>41</sub> is Gly or Ala; X<sub>42</sub> is Gly or Ala; X<sub>43</sub> is Trp or Ala; X<sub>44</sub> is Asn or Ala; X<sub>45</sub> is Tyr or Trp; X<sub>46</sub> is Met or Gln;

(viii) FSGYYWS (SEQ ID NO:8);

(ix) EINHSGSTNYKTSLS (SEQ ID NO:9); or

(x) GLLRGGWNDVDYXXGMDV (SEQ ID NO:10), or

3) a peptide comprising consecutive amino acids having a sequence of any of a) and b), or at least two of the sequences in (a)(i), (a)(ii) and (b)(i) through (b)(x), or

4) a peptide comprising consecutive amino acids having a sequence comprising at least two identical sequences included in (a)(i), (a)(ii) and (b)(i) through (b)(x);

c) adjusting the pH of the solution of step b) until the peptide dissolves in the solution; and

d) if necessary, adjusting the pH of the solution of step c) to a pH of 4-9, thereby manufacturing the pharmaceutical composition.

Claims 17-23. (Canceled)

24. (Previously Presented) A composition prepared by the process of claim 16.

25. (Original) A lyophilized pharmaceutical composition comprising from 0.1 mg/ml to 20 mg/ml of the composition of a pharmaceutically acceptable salt of

a) a peptide comprising at least 12 and at most 30

consecutive amino acids having a sequence corresponding to

- (i) a sequence of amino acids found within a complementarity-determining region (CDR) of a heavy or a light chain of a human monoclonal anti-DNA 16/6 Id antibody, or
  - (ii) a sequence of amino acids found within a complementarity-determining region (CDR) of a heavy or a light chain of a pathogenic anti-DNA monoclonal antibody that induces a systemic lupus erythematosus (SLE)-like disease response in mice, or
- b) a peptide comprising consecutive amino acids having the sequence
- (i) TGYXX<sub>1</sub>X<sub>2</sub>X<sub>3</sub>X<sub>4</sub>X<sub>5</sub>QSPEKSLEWIG (SEQ ID NO:11)  
wherein X<sub>1</sub> is Met, Ala or Val; X<sub>2</sub> is Gln, Asp, Glu or Arg; X<sub>3</sub> is Trp or Ala; X<sub>4</sub> is Val or Ser; and X<sub>5</sub> is Lys, Glu or Ala;
  - (ii) EINPSTGGX<sub>6</sub>X<sub>7</sub>X<sub>8</sub>X<sub>9</sub>X<sub>10</sub>X<sub>11</sub>X<sub>12</sub>KAKAT (SEQ ID NO:12)  
wherein X<sub>6</sub> and X<sub>7</sub> are each Thr, Val or Ala; X<sub>8</sub> is Tyr or Phe; X<sub>9</sub> is Asn or Asp; X<sub>10</sub> is Gln or Glu; X<sub>11</sub> is Lys or Glu, and X<sub>12</sub> is Phe or Tyr;
  - (iii) YYCARX<sub>13</sub>X<sub>14</sub>X<sub>15</sub>X<sub>16</sub>PYAX<sub>17</sub>X<sub>18</sub>YWGQGS (SEQ ID NO:13)  
wherein X<sub>13</sub> is Phe, Thr or Gly; X<sub>14</sub> is Leu, Ala or Ser; X<sub>15</sub> is Trp or Ala; X<sub>16</sub> is Glu or Lys; X<sub>17</sub> is Met or Ala, and X<sub>18</sub> is Asp, Lys or Ser;
  - (iv) GYNX<sub>19</sub>X<sub>20</sub>X<sub>21</sub>X<sub>22</sub>X<sub>23</sub>X<sub>24</sub>SHGX<sub>25</sub>X<sub>26</sub>LEWIG (SEQ ID NO:14)  
wherein X<sub>19</sub> is Met or Ala; X<sub>20</sub> is Asn, Asp or Arg; X<sub>21</sub> is Trp or Ala; X<sub>22</sub> is Val or Ser; X<sub>23</sub> is Lys or Glu; X<sub>24</sub> is Gln or Ala; X<sub>25</sub> is Lys or Glu, and X<sub>26</sub> is Ser or Ala;
  - (v) YYCARX<sub>27</sub>X<sub>28</sub>X<sub>29</sub>YGX<sub>30</sub>X<sub>31</sub>X<sub>32</sub>GQTL (SEQ ID NO:15)  
wherein X<sub>27</sub> is Ser or Phe; X<sub>28</sub> is Gly or Ala; X<sub>29</sub> is



Arg, Ala or Glu; X<sub>30</sub> is Asn or Asp; X<sub>31</sub> is Tyr or Phe, and X<sub>32</sub> is Trp, His or Ala;

(vi) X<sub>33</sub>YYWSWIX<sub>34</sub>QX<sub>35</sub>PX<sub>36</sub>X<sub>37</sub>GX<sub>38</sub>EWIG (SEQ ID NO:16)

wherein X<sub>33</sub> is Gly or Thr Gly; X<sub>34</sub> is Arg or Lys; X<sub>35</sub> is Pro or Ser; X<sub>36</sub> is Gly or Glu; X<sub>37</sub> is Lys or Asp; and X<sub>38</sub> is Glu, Leu or Ser;

(vii) YYCARX<sub>39</sub>LLX<sub>40</sub>X<sub>41</sub>X<sub>42</sub>X<sub>43</sub>X<sub>44</sub>DVDYX<sub>45</sub>GX<sub>46</sub>DV (SEQ ID NO:17)

wherein X<sub>39</sub> is Gly or Phe; X<sub>40</sub> is Arg or Ala; X<sub>41</sub> is Gly or Ala; X<sub>42</sub> is Gly or Ala; X<sub>43</sub> is Trp or Ala; X<sub>44</sub> is Asn or Ala; X<sub>45</sub> is Tyr or Trp; X<sub>46</sub> is Met or Gln;

(viii) FSGYYWS (SEQ ID NO:8);

(ix) EINHSGSTNYKTSLS (SEQ ID NO:9); or

(x) GLLRGGWNDVDYGGMDV (SEQ ID NO:10), or

c) a peptide comprising consecutive amino acids having a sequence of any of a) and b), or at least two of the sequences in (a)(i), (a)(ii) and (b)(i) through (b)(x), or

d) a peptide comprising consecutive amino acids having a sequence comprising at least two identical sequences included in (a)(i), (a)(ii) and (b)(i) through (b)(x); and

a solubility enhancer selected from the group consisting of dimethyl-acetamide, polyethylene glycol, polyoxylated castor oil, N-methyl-2-pyrrolidinone, 1-ethenyl-2-pyrrolidinone, polyoxyethylene sorbitan esters, and a substituted  $\beta$ -cyclodextrin.

26. (Canceled)

27. (Withdrawn-Currently Amended) A process ~~of lyophilizing for~~ manufacturing the lyophilized pharmaceutical composition of claim ~~425~~, comprising the steps of:

a) preparing a solution of dimethyl-acetamide, polyethylene glycol, polyoxylated castor oil, N-methyl-2-pyrrolidinone, 1-ethenyl-2-pyrrolidinone, polyoxyethylene sorbitan esters, or a substituted  $\beta$ -cyclodextrin in an aqueous carrier at a predetermined concentration;

b) adding a predetermined amount of a pharmaceutically acceptable salt of

1) a peptide comprising at least 12 and at most 30 consecutive amino acids having a sequence corresponding to

(i) a sequence of amino acids found within a complementarity-determining region (CDR) of a heavy or a light chain of a human monoclonal anti-DNA 16/6 Id antibody, or

(ii) a sequence of amino acids found within a complementarity-determining region (CDR) of a heavy or a light chain of a pathogenic anti-DNA monoclonal antibody that induces a systemic lupus erythematosus (SLE)-like disease response in mice,

2) a peptide comprising amino acids having the sequence

(i) TGYX<sub>1</sub>X<sub>2</sub>X<sub>3</sub>X<sub>4</sub>X<sub>5</sub>QSPEKSLEWIG (SEQ ID NO:11)

wherein X<sub>1</sub> is Met, Ala or Val; X<sub>2</sub> is Gln, Asp, Glu or Arg; X<sub>3</sub> is Trp or Ala; X<sub>4</sub> is Val or Ser; and X<sub>5</sub> is Lys, Glu or Ala;

(ii) EINPSTGGX<sub>6</sub>X<sub>7</sub>X<sub>8</sub>X<sub>9</sub>X<sub>10</sub>X<sub>11</sub>X<sub>12</sub>KAKAT (SEQ ID NO:12)

wherein X<sub>6</sub> and X<sub>7</sub> are each Thr, Val or Ala; X<sub>8</sub> is Tyr or Phe; X<sub>9</sub> is Asn or Asp; X<sub>10</sub> is Gln or Glu; X<sub>11</sub> is Lys or Glu, and X<sub>12</sub> is Phe or Tyr;

(iii) YYCARX<sub>13</sub>X<sub>14</sub>X<sub>15</sub>X<sub>16</sub>PYAX<sub>17</sub>X<sub>18</sub>YWGQGS (SEQ ID NO:13)

wherein X<sub>13</sub> is Phe, Thr or Gly; X<sub>14</sub> is Leu, Ala or Ser; X<sub>15</sub> is Trp or Ala; X<sub>16</sub> is Glu or Lys; X<sub>17</sub> is Met or Ala, and X<sub>18</sub> is Asp, Lys or Ser;

(iv) GYNX<sub>19</sub>X<sub>20</sub>X<sub>21</sub>X<sub>22</sub>X<sub>23</sub>X<sub>24</sub>SHGX<sub>25</sub>X<sub>26</sub>LEWIG (SEQ ID NO:14)

wherein X<sub>19</sub> is Met or Ala; X<sub>20</sub> is Asn, Asp or Arg;  
X<sub>21</sub> is Trp or Ala; X<sub>22</sub> is Val or Ser; X<sub>23</sub> is Lys or  
Glu; X<sub>24</sub> is Gln or Ala; X<sub>25</sub> is Lys or Glu, and X<sub>26</sub> is  
Ser or Ala;

(v) YYCARX<sub>27</sub>X<sub>28</sub>X<sub>29</sub>YGX<sub>30</sub>X<sub>31</sub>X<sub>32</sub>GQTL (SEQ ID NO:15)

wherein X<sub>27</sub> is Ser or Phe; X<sub>28</sub> is Gly or Ala; X<sub>29</sub> is  
Arg, Ala or Glu; X<sub>30</sub> is Asn or Asp; X<sub>31</sub> is Tyr or  
Phe, and X<sub>32</sub> is Trp, His or Ala;

(vi) X<sub>33</sub>YYWSWIX<sub>34</sub>QX<sub>35</sub>PX<sub>36</sub>X<sub>37</sub>GX<sub>38</sub>EWIG (SEQ ID NO:16)

wherein X<sub>33</sub> is Gly or Thr Gly; X<sub>34</sub> is Arg or Lys; X<sub>35</sub>  
is Pro or Ser; X<sub>36</sub> is Gly or Glu; X<sub>37</sub> is Lys or Asp;  
and X<sub>38</sub> is Glu, Leu or Ser;

(vii) YYCARX<sub>39</sub>LLX<sub>40</sub>X<sub>41</sub>X<sub>42</sub>X<sub>43</sub>X<sub>44</sub>DVDYX<sub>45</sub>GX<sub>46</sub>DV (SEQ ID NO:17)

wherein X<sub>39</sub> is Gly or Phe; X<sub>40</sub> is Arg or Ala; X<sub>41</sub> is  
Gly or Ala; X<sub>42</sub> is Gly or Ala; X<sub>43</sub> is Trp or Ala; X<sub>44</sub>  
is Asn or Ala; X<sub>45</sub> is Tyr or Trp; X<sub>46</sub> is Met or Gln;

(viii) FSGYYWS (SEQ ID NO:8);

(ix) EINHSGSTNYKTSLS (SEQ ID NO:9); or

(x) GLLRGGWNDVDYYYGMDV (SEQ ID NO:10), or

3) a peptide comprising consecutive amino acids having  
a sequence of any of a) and b), or at least two of the  
sequences in (a)(i), (a)(ii) and (b)(i) through (b)(x),  
or

4) a peptide comprising consecutive amino acids having  
a sequence comprising at least two identical sequences  
included in (a)(i), (a)(ii) and (b)(i) through (b)(x);

c) adjusting the pH of the solution of step b) until the  
peptide dissolves in the solution;

d) if necessary, adjusting the pH of the solution of step c)  
to a pH of 4-9, thereby manufacturing the pharmaceutical  
composition; and

e) lyophilizing the pharmaceutical composition of step d) by:

a-i) lowering the temperature of the pharmaceutical composition to -40°C;  
ba-ii) holding the temperature at -40°C for a predetermined time;  
ea-iii) raising the temperature of the solution to 20°C;  
da-iv) holding the temperature at 20°C for a predetermined time; and  
ea-v) reducing the pressure and holding the temperature at 20°C for a predetermined time, thereby lyophilizing the pharmaceutical composition;

or-

b-i) lowering the temperature of the pharmaceutical composition to -45°C;  
b-ii) holding the temperature at -45°C for a predetermined time;  
b-iii) raising the temperature of the solution to -20°C;  
b-iv) raising the temperature of the solution to 25°C; and  
b-v) holding the temperature at 25°C for a predetermined time, thereby lyophilizing the pharmaceutical composition.

Claims 28-35. (Canceled)

36. (Withdrawn-Currently Amended) The process of claim 27, wherein  
step a-i) is performed within 2 hours;  
step ba-ii) is performed within 3 hours;  
step ea-iii) is performed over 13 hours and at a pressure of 110µbar;  
step da-iv) is performed over 13 hours and at a pressure of 110µbar; and  
step ea-v) is performed over 5 hours and the pressure is reduced to 10µbar.

Applicants: Sharon Cohen-Vered et al.  
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37. (Withdrawn) A lyophilized pharmaceutical composition prepared by the process of claim 27.

Claims 38-46. (Canceled)

47. (Withdrawn-Currently Amended) The process of claim ~~38~~27, wherein

step ~~a~~b-i) is performed within 6 hours;  
step ~~b~~b-ii) is performed within 3 hours;  
step ~~c~~c-iii) is performed over 19 hours and at a pressure of 150μbar;  
step ~~d~~d-iv) is performed over 13 hours and at a pressure of 150μbar; and  
step ~~e~~e-v) is performed over 8 hours and at a pressure of 150μbar.

Claims 48-51. (Canceled)

52. (Currently Amended) A packaged pharmaceutical composition comprised of:  
a packaging material; and  
a predetermined amount of the lyophilized pharmaceutical composition of claim 37-~~or 48~~.